

CRF Errors Corrected by the STIC Systems Branch

01PE 02 CO

Serial Number: _____

09/369,236

CRF Processing Date: _____

8/24/99

Edited by: _____

Verified by: TC (STIC staff)

- ☐ Changed a file from non-ASCII to ASCII
- ☐ Changed the margins in cases where the sequence text was "wrapped" down to the next line.
- ☐ Edited a format error in the Current Application Data section, specifically: _____
- ☐ Edited the Current Application Data section with the actual current number. The number inputted by the applicant was ☐ the prior application data; or ☐ other ENTERED.
- ☐ Added the mandatory heading and subheadings for "Current Application Data".
- ☐ Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.
- ☐ Changed the spelling of a mandatory field (the headings or subheadings), specifically: _____
- ☐ Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: _____
- ☐ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: _____
- ☒ Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.
- ☐ Inserted colons after headings/subheadings. Headings edited included: _____
- ☒ Deleted extra, invalid, headings used by an applicant, specifically: (A) NAME!
- ☐ Deleted: ☐ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end of file; ☐ page numbers throughout text; ☐ other invalid text, such as _____
- ☐ Inserted mandatory headings, specifically: _____
- ☐ Corrected an obvious error in the response, specifically: _____
- ☐ Edited identifiers where upper case is used but lower case is required, or vice versa.
- ☐ Corrected an error in the Number of Sequences field, specifically: _____
- ☐ A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
- ☐ Deleted *ending* stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: _____
- ☐ Other: _____

*Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form. 3/1/95

RAW SEQUENCE LISTING
PATENT APPLICATION US/09/369,236

DATE: 08/24/1999
TIME: 14:38:53

INPUT SET: S33041.raw

This Raw Listing contains the General
Information Section and up to the first 5 pages.

SEQUENCE LISTING

Does Not Comply
Corrected Diskette Needed

(1) General Information:

(i) APPLICANT: ↑

~~(A) NAME:~~ Grant A. Krafft, William L. Klein, Brett A. Chromy,
Mary P. Lambert, Caleb E. Finch, Todd Morgan, Pat Wals,
Irina Rozovsky, Ann Barlow

(ii) TITLE OF INVENTION: Amyloid (Protein (Globular Assembly
and Uses Thereof)

(iii) NUMBER OF SEQUENCES: 4

(iv) CORRESPONDENCE ADDRESS:

(A) ADDRESSEE: McDonnell Boehnen Hulbert & Berghoff
(B) STREET: 300 South Wacker Drive
(C) CITY: Chicago
(D) STATE: IL
(E) COUNTRY: USA
(F) ZIP: 60606

(v) COMPUTER READABLE FORM:

(A) MEDIUM TYPE: Floppy disk
(B) COMPUTER: IBM PC compatible
(C) OPERATING SYSTEM: PC-DOS/MS-DOS
(D) SOFTWARE: PatentIn Release #1.0, Version #1.30 (US)

(vi) CURRENT APPLICATION DATA:

(A) APPLICATION NUMBER: unassigned
(B) FILING DATE: 04-AUG-1999
(C) CLASSIFICATION DATA:

(vii) PRIOR APPLICATION DATA:

(A) APPLICATION NUMBER: US 08/796,089
(B) FILING DATE: 05-FEB-1997

(2) INFORMATION FOR SEQ ID NO: 1:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 20 base pairs
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

-->
-->

--> OK

RAW SEQUENCE LISTING
PATENT APPLICATION US/09/369,236DATE: 08/24/1999
TIME: 14:38:53

INPUT SET: S33041.raw

47 (ii) MOLECULE TYPE: other nucleic acid
48
49
50 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:
51
52 GCACCTTCTT TCCCTTCATC20
53
54 (2) INFORMATION FOR SEQ ID NO: 2:
55
56 (i) SEQUENCE CHARACTERISTICS:
57 (A) LENGTH: 20 base pairs
58 (B) TYPE: nucleic acid
59 (C) STRANDEDNESS: single
60 (D) TOPOLOGY: linear
61
62 (ii) MOLECULE TYPE: other nucleic acid
63
64
65 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2:
66
67 TGCTGATGTA CCAGTTGGGG20
68
69
70 (2) INFORMATION FOR SEQ ID NO: 3:
71
72 (i) SEQUENCE CHARACTERISTICS:
73 (A) LENGTH: 19 base pairs
74 (B) TYPE: nucleic acid
75 (C) STRANDEDNESS: single
76 (D) TOPOLOGY: linear
77
78 (ii) MOLECULE TYPE: other nucleic acid
79
80
81 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 3:
82
83 CAGTCCTTGA CCTGCGACC 19
84
85
86 (2) INFORMATION FOR SEQ ID NO: 4:
87
88 (i) SEQUENCE CHARACTERISTICS:
89 (A) LENGTH: 19 base pairs
90 (B) TYPE: nucleic acid
91 (C) STRANDEDNESS: single
92 (D) TOPOLOGY: linear
93
94 (ii) MOLECULE TYPE: other nucleic acid
95
96 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 4:
97
98 GCCTCACATC ACATCCTTG 19
99

SEQUENCE VERIFICATION REPORT
PATENT APPLICATION US/09/369,236DATE: 08/24/1999
TIME: 14:38:54**INPUT SET: S33041.raw**

Line	Error	Original Text
5	Mandatory Value Not Present	(i) APPLICANT:
6	Unknown or Misplaced Identifier	(A) NAME: Grant A. Krafft, William L. Klein, Brett A. Ch
30	Wrong application Serial Number	(A) APPLICATION NUMBER: unassigned